

Amendments to the Specification:

Please insert the following new paragraph entitled: **CROSS REFERENCE TO RELATED APPLICATIONS** before "**STATEMENTS AS TO RIGHTS TO INVENTIONS MADE UNDER FEDERALLY SPONSORED RESEARCH AND DEVELOPMENT**".

--CROSS REFERENCE TO RELATED APPLICATIONS

This application was filed under 35 U.S.C. 371 as a National stage application from Serial No. PCT/US00/22226, filed August 11, 2000, which claims the benefit of Provisional Patent Application Serial No. 60/148,907, filed August 13, 1999, the disclosures of each are incorporated herein by reference.--

BRIEF DESCRIPTION OF THE DRAWINGS:

Please replace the paragraph entitled Figure 4A with the following amended paragraph:

Figure [[4A]] 4. Exposed hydrophobic residues on the factor VIII C2 domain.

The orientation is the same as Figure 3. The protein displays two distinct exposed hydrophobic surfaces. The first, at the upper end of the β -sandwich, includes Phe 2275, Tyr 2332 and Leu 2302. The second surface, formed by two β -turns and a loop as described in Figure 3, includes Met 2199 and Phe 2200 from the first turn, Leu 2251 and 2252 from the second turn, and Val 2223 from the loop. As shown in Figure 5, these structures extend approximately 10 Å beyond the protein core and flank a pair of positively charged clefts. This structure therefore appears optimal for associating with negatively charged phospholipid membranes. Figure 4B: The In the right side of Figure 4 the protein is rotated clockwise by approximately 45° relative to panel a the left side of Figure 4, in order to place the hydrophobic residues (Met 2199, Phe 2200, Leu 2151, Leu 2152, and Val 2223) and underlying basic residues (Arg 2215, Arg 2220, Lys 2249 and Lys 2227) along a horizontal axis (grey line) that represents the predicted position of the polar/nonpolar boundary of the phospholipid bilayer.